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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/070,503

DATE: 11/19/2002 P.6
TIME: 16:52:14

Input Set : **A:\10.070503.txt**
Output Set: **N:\CRF4\11192002\J070503.raw**

```
 3 <110> APPLICANT: Viventia Biotech Inc.
 5 <120> TITLE OF INVENTION: ENHANCED PHAGE DISPLAY LIBRARIES AND METHODS FOR
 6       PRODUCING SAME
 8 <130> FILE REFERENCE: 33956-49
```
C--> `10 <140> CURRENT APPLICATION NUMBER: US/10/070,503`
```
11 <141> CURRENT FILING DATE: 2000-09-07
13 <150> PRIOR APPLICATION NUMBER: PCT/CA00/01027
14 <151> PRIOR FILING DATE: 2000-09-07
16 <150> PRIOR APPLICATION NUMBER: CA2282179
17 <151> PRIOR FILING DATE: 1999-09-07
19 <150> PRIOR APPLICATION NUMBER: US60/163,546
20 <151> PRIOR FILING DATE: 1999-11-04
22 <160> NUMBER OF SEQ ID NOS: 60
24 <170> SOFTWARE: PatentIn Ver. 2.1
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 396
28 <212> TYPE: DNA
29 <213> ORGANISM: human
31 <400> SEQUENCE: 1
32 gaggtccagc tgcaggagtc tggggggaggc ttagtccagc ctggggggtc cctgagactc 60
33 tcctgttcag cctctggatt caccttcagt agctatgcta tgcactgggt ccgccaggct 120
34 ccagggaagg gactggaata tgtttcagct attagtagta atggggggtag cacatactac 180
35 gcagactccg tgaagggcag attcaccatc tccagagaca attccaagaa cactctgtat 240
36 cttcaaatga gcagtctgag agctgaggac acggctgtgt attactgtgt gaaagacagg 300
37 ttaaaagtgg agtactatga tagtagtggt tattacgttt ctcggttcgg tgctttttgat 360
38 atctggggcc aagggacaac ggtcaccgtc tcatca                           396
41 <210> SEQ ID NO: 2
42 <211> LENGTH: 132
43 <212> TYPE: PRT
44 <213> ORGANISM: human
46 <400> SEQUENCE: 2
47 Glu Val Gln Leu Gln Glu Ser Gly Gly Gly Leu Val Gln Pro Gly Gly
48  1               5                  10                  15
50 Ser Leu Arg Leu Ser Cys Ser Ala Ser Gly Phe Thr Phe Ser Ser Tyr
51              20                  25                  30
53 Ala Met His Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Tyr Val
54          35                  40                  45
56 Ser Ala Ile Ser Ser Asn Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val
57      50                  55                  60
59 Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr
60 65                  70                  75                  80
62 Leu Gln Met Ser Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys
63              85                  90                  95
```

```
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              Input Set : A:\10.070503.txt
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  65 Val Lys Asp Arg Leu Lys Val Glu Tyr Tyr Asp Ser Ser Gly Tyr Tyr
  66             100             105             110
  68 Val Ser Arg Phe Gly Ala Phe Asp Ile Trp Gly Gln Gly Thr Thr Val
  69         115             120             125
  71 Thr Val Ser Ser
  72     130
  75 <210> SEQ ID NO: 3
  76 <211> LENGTH: 5
  77 <212> TYPE: PRT
  78 <213> ORGANISM: human
  80 <400> SEQUENCE: 3
  81 Ser Tyr Ala Met His
  82   1               5
  85 <210> SEQ ID NO: 4
  86 <211> LENGTH: 16
  87 <212> TYPE: PRT
  88 <213> ORGANISM: human
  90 <400> SEQUENCE: 4
  91 Ala Ile Ser Ser Asn Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val Lys
  92   1               5               10              15
  95 <210> SEQ ID NO: 5
  96 <211> LENGTH: 23
  97 <212> TYPE: PRT
  98 <213> ORGANISM: human
 100 <400> SEQUENCE: 5
 101 Asp Arg Leu Lys Val Glu Tyr Tyr Asp Ser Ser Gly Tyr Tyr Val Ser
 102   1               5               10              15
 104 Arg Phe Gly Ala Phe Asp Ile
 105                 20
 108 <210> SEQ ID NO: 6
 109 <211> LENGTH: 33
 110 <212> TYPE: DNA
 111 <213> ORGANISM: Artificial Sequence
 113 <220> FEATURE:
 114 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
 116 <220> FEATURE:
W--> 117 <221> NAME/KEY: variable residue
 118 <222> LOCATION: (16)
 119 <223> OTHER INFORMATION: The nucleotide at this position may be a or c
 121 <220> FEATURE:
W--> 122 <221> NAME/KEY: variable residue
 123 <222> LOCATION: (17)
 124 <223> OTHER INFORMATION: The nucleotide at this position may be a, c, t or g
 126 <220> FEATURE:
W--> 127 <221> NAME/KEY: variable residue
 128 <222> LOCATION: (18)
 129 <223> OTHER INFORMATION: The nucleotide at this position may be a, c, t or g
 131 <220> FEATURE:
W--> 132 <221> NAME/KEY: variable length
```

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```
     133 <222> LOCATION: (16)...(18)
     134 <223> OTHER INFORMATION: The nucleotides at these positions may be repeated
     136 <400> SEQUENCE: 6
W--> 137 gccccagata tcaaannntt tcacacagta ata                          33
     140 <210> SEQ ID NO: 7
     141 <211> LENGTH: 18
     142 <212> TYPE: DNA
     143 <213> ORGANISM: Artificial Sequence
     145 <220> FEATURE:
     146 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
     148 <400> SEQUENCE: 7
     149 tgttcagcta gcggattc                                           18
     152 <210> SEQ ID NO: 8
     153 <211> LENGTH: 42
     154 <212> TYPE: DNA
     155 <213> ORGANISM: Artificial Sequence
     157 <220> FEATURE:
     158 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
     160 <400> SEQUENCE: 8
     161 tgaggagacg gtgaccgttg tcccttggcc ccagatatca aa                42
     164 <210> SEQ ID NO: 9
     165 <211> LENGTH: 38
     166 <212> TYPE: DNA
     167 <213> ORGANISM: Artificial Sequence
     169 <220> FEATURE:
     170 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
     172 <400> SEQUENCE: 9
     173 catgaccaca gtgcacagga ggtccagctg caggagtc                     38
     176 <210> SEQ ID NO: 10
     177 <211> LENGTH: 18
     178 <212> TYPE: DNA
     179 <213> ORGANISM: Artificial Sequence
     181 <220> FEATURE:
     182 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
     184 <400> SEQUENCE: 10
     185 tttcacacag taatacac                                           18
     188 <210> SEQ ID NO: 11
     189 <211> LENGTH: 57
     190 <212> TYPE: DNA
     191 <213> ORGANISM: Artificial Sequence
     193 <220> FEATURE:
     194 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
     196 <400> SEQUENCE: 11
     197 cgattctgcg gccgctgagg agacggtgac cgttgtccct tggccccaga tatcaaa  57
     200 <210> SEQ ID NO: 12
     201 <211> LENGTH: 35
     202 <212> TYPE: DNA
     203 <213> ORGANISM: Artificial Sequence
     205 <220> FEATURE:
```

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```
      206 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      208 <220> FEATURE:
W-->  209 <221> NAME/KEY: variable residue
      210 <222> LOCATION: (18)
      211 <223> OTHER INFORMATION: The nucleotide at this position may be a or c
      213 <220> FEATURE:
W-->  214 <221> NAME/KEY: variable residue
      215 <222> LOCATION: (19)
      216 <223> OTHER INFORMATION: The nucleotide at this position may be a, g, t or c
      218 <220> FEATURE:
W-->  219 <221> NAME/KEY: variable residue
      220 <222> LOCATION: (20)
      221 <223> OTHER INFORMATION: The nucleotide at this position may be a, g, t or c
      223 <220> FEATURE:
W-->  224 <221> NAME/KEY: variable length
      225 <222> LOCATION: (18)...(20)
      226 <223> OTHER INFORMATION: The nucleotides at these positions may be repeated
      228 <400> SEQUENCE: 12
W-->  229 gttgtccctt ggccccannn tttcacacag taata                      35
      232 <210> SEQ ID NO: 13
      233 <211> LENGTH: 30
      234 <212> TYPE: DNA
      235 <213> ORGANISM: Artificial Sequence
      237 <220> FEATURE:
      238 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      240 <400> SEQUENCE: 13
      241 actttcttgt aattggacct cggcctgcgc                            30
      244 <210> SEQ ID NO: 14
      245 <211> LENGTH: 21
      246 <212> TYPE: DNA
      247 <213> ORGANISM: Artificial Sequence
      249 <220> FEATURE:
      250 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      252 <400> SEQUENCE: 14
      253 ctctcctgtg ctgcctctgg a                                     21
      256 <210> SEQ ID NO: 15
      257 <211> LENGTH: 21
      258 <212> TYPE: DNA
      259 <213> ORGANISM: Artificial Sequence
      261 <220> FEATURE:
      262 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      264 <400> SEQUENCE: 15
      265 tccagaggca gcacaggaga g                                     21
      268 <210> SEQ ID NO: 16
      269 <211> LENGTH: 54
      270 <212> TYPE: DNA
      271 <213> ORGANISM: Artificial Sequence
      273 <220> FEATURE:
      274 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
```

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                Input Set : A:\10.070503.txt
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      276 <400> SEQUENCE: 16
      277 cgcacagtaa tacacagccg tgtcctcagc tctcagactg ttcatttgaa gata   54
      280 <210> SEQ ID NO: 17
      281 <211> LENGTH: 24
      282 <212> TYPE: DNA
      283 <213> ORGANISM: Artificial Sequence
      285 <220> FEATURE:
      286 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      288 <400> SEQUENCE: 17
      289 gtgtattact gtgcgaaaga cagg   24
      292 <210> SEQ ID NO: 18
      293 <211> LENGTH: 21
      294 <212> TYPE: DNA
      295 <213> ORGANISM: Artificial Sequence
      297 <220> FEATURE:
      298 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      300 <400> SEQUENCE: 18
      301 caattacaag ctagtggtgg c   21
      304 <210> SEQ ID NO: 19
      305 <211> LENGTH: 39
      306 <212> TYPE: DNA
      307 <213> ORGANISM: Artificial Sequence
      309 <220> FEATURE:
      310 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      312 <400> SEQUENCE: 19
      313 tatggatcct gaggagacgg tgacctgtgt cccttggcc   39
      316 <210> SEQ ID NO: 20
      317 <211> LENGTH: 38
      318 <212> TYPE: DNA
      319 <213> ORGANISM: Artificial Sequence
      321 <220> FEATURE:
      322 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      324 <400> SEQUENCE: 20
      325 catgaccaca gtgcacagga ggtccaatta caagaaag   38
      328 <210> SEQ ID NO: 21
      329 <211> LENGTH: 42
      330 <212> TYPE: DNA
      331 <213> ORGANISM: Artificial Sequence
      333 <220> FEATURE:
      334 <223> OTHER INFORMATION: Description of Artificial Sequence:primer
      336 <220> FEATURE:
W--> 337 <221> NAME/KEY: variable residue
      338 <222> LOCATION: (22)
      339 <223> OTHER INFORMATION: The nucleotide at this position may be a or c
      341 <220> FEATURE:
W--> 342 <221> NAME/KEY: variable residue
      343 <222> LOCATION: (23)
      344 <223> OTHER INFORMATION: The nucleotide at this position may be a, g, t or c
      346 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 11/19/2002
PATENT APPLICATION: **US/10/070,503**   TIME: 16:52:15

Input Set : **A:\10.070503.txt**
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<u>Please Note:</u>
**Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.**

Seq#:6; N Pos. 16,17,18
Seq#:12; N Pos. 18,19,20
Seq#:21; N Pos. 22,23,24

VERIFICATION SUMMARY   DATE: 11/19/2002
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```
L:10  M:270 C: Current Application Number differs, Replaced Current Application Number
L:117 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:122 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:127 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:132 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:6
L:137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:209 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:12
L:214 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:12
L:219 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:12
L:224 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:12
L:229 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:337 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:21
L:342 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:21
L:347 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:21
L:352 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:21
L:357 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
```